## AMENDMENTS TO THE DRAWINGS

Replace the original Fig. 1A and 1B with the attached sheet of drawings at Tab 1.

The following is an explanation of the changes made to the immediate prior version of the Figures: Pending Fig. 1A. has been amended to correct a typographical error by changing "G" to "H" as shown in red ink in Tab 2. Pending Fig. 1B. has been amended to correct a typographical error by changing "F" to "G" as shown in red ink in Tab 2. Support is in the originally filed Figure 1A and 1B (Tab 3) of the priority provisional Application No. 60/081,734 (our File UCSD-04729).

Enclosure:

Replacement Sheets (Tab 1)

Annotated Sheet Showing Changes In Red Ink (Tab 2)

Originally filed Sheets in Application No. 60/081,734 (Tab 3)

1 MSVDEICENTKMGREYALLGNYETSLVYYQGVLQQIQKLL 41 TSVHEPQRKHQWQTIRQELSQEYEHVKNITKTLNGFKSEP 81 AAPEPAPNHRAAPFSHHQHAAKPAAAEPARDPDVWPPPTP 121 **VDHRPSPPYQRAARKDPPRRSEPSKPANRAPGNDRGGRGP** 161 SDRRGDARSGGGGRGGARGSDKDKNRGGKSDKDKKAPSGE 201 **EGDEKKFDPAGYDKDLVENLERDIVQRNPNVHWADIAGLT** 241 <u>EAKRLLEEAVVLPLWMPDYFKGIRRPWKGVLMVGPPGTGK</u> 281  $oldsymbol{\Pi}$ MLAKAVATECGTTFFNVSSASLTSKYHGESEKLVRLLFE 321 MARFYAPSTIFIDEIDSICSKRGTGSEHEASRRVKSELLI 361 QMDGVSGPSAGEESSKMVMVLAATNFPWDIDEALRRRLEK 401 RIYIPLPEIDGREQLLRINLKEVPLADDIDLKSIAEKMDG 441 YSGADITNVCRDASMMAMRRRIQGLRPEEIRHIPKEELNQ 481 **PSTPADFLLALQKVSKSVGKEDLVKYMAWMEEFGSV** 

## FIG. 1A

```
VHWADIAGLTEAKRLLEEAVVLPLWMPDYEKGIRRPW.KGVLMVGPPGTG
MSLDDIIGMHDVKOVLHEAVTLPLLVPEFFOGLRSPW.KAMVLAGPPGTG
STYDMVGGLTKOIKEIKEVIELPVKHPELFESLGIAOPKGVILYGPPGTG
TTFADVAGCDEAKEEVAELVEY.LREPSRFOKLGGKIPKGVLMVGPPGTG
IKWGDIGALANAKOVLLETLEWPTKYEPIEVNCCPLRISGILLYGYPGCG
p60
                    190
mei-l
Sug1p
                    145
                    149
ftsH
                                IKWGDIGALANAKOVLLETLEWPTKYEPIFVNCPLRLRSGILLYGYPGCGEKMGIGGLDDEFSDIFRRAFASRVFPPEIVEQMGCKHVKGILLYGPPGCG
Paslp
                   694
NSF
                    224
                               KTMLAKAVATECGTTEFN. VSSASLTSKYHGESEKLVRILFEMARFYAP.
KTLIARAIASESSSTEFT. VSSTDLSSKWRGDSEKIVRILFELARFYAP.
KTLLARAVAHHTDCKFIR. VSGAELVOKYIGEGSRMVRELFVMAREHAP.
KTLLAKAIAGEAKVPEFT. ISGSDFVEMFVGVGASRVRDMFEOAKKAAP.
KTLLASAVAOOCGLNEIS. VKGPEILNKFIGASEONIRELFERAOSVKP.
KTLLAROIGKMLNAREPKVVNGPEILNKYMGESEANIRKLFADAEEEQRR
p60
mei-1
                    239
                    195
Sug1p
                    198
ftsH
Paslp
                    744
                    274
NSF
                               ....STIFIDEIDSICSKR..GTGSEHEASRRVKSELLIOMDGVSGP
....STIFIDEIDTLGGOR..GNSGEHEASRRVKSEFLVOMDG....
STIFMDEIDSIGSTRVEGSGGDSEVORTMLELLNOLDGFET.
....CTIFIDEIDAVGRORGAGLGGGHDEREOTLNOMLVEMDGFEG.
....CTLEFDEFDSIAPKRGH...DSTGVTDRVVNOLLTOMDGAEGL
LGANSGLHTITFDEIDAICKORGS.MAGSTGVHDTVVNOLLSKIDGVEQL
p60
                    328
                    287
243
mei−1
Sug1p
                    246
792
ftsH
Pas1p
                    324
NSF
                               SAGEESSKMVMVLAATNFPWDIDEALRR..RLEKRIYIPLPEIDGREOLL
SONKFDSRRVFVLAATNIPWELDEALRR..RFEKRIFIPLPDIDARKKLI
....SKNIKIMATNRLDILDPALLRPGRIDRKIEFPPPSVAARAEIL
....NEGIIVIAATNRPDVLDPALLRPGRFDROVVVCLPDVRGREOIL
D.....GVYILAATSRPDLIDSALLRPGRLDKSVICNIPTESERLDIL
                   369
324
285
288
p60
mei-l
Sug1p
ftsH
Paslp
                                N.....NILVIGMTNRPDLIDEALLRPGRLEVKMEIGLPDEKGRLOIL
NSF
                                RINL.....KEVPLADDIDLKSIAEKMDGYSGADIT
EKSM....EGTPKSDEINYDDLAARTEGFSGADVV
RIHS....RKMNLTRGINLRKVAEKMNGCSGADVK
KVHM....RRVPLAPDIDAAIIARGTPGFSGADLA
QAIVNSKDKDTGQKKFALEKNADLKLIAEKTAGFSGADLQ
HI...HTARMRGHQ..LLSADVDIKELAVETKNFSGAELE
p60
mei-1
                    372
                   329
332
875
Sug1p
 ftšH
Paslp
NSF
```

FIG. 1B

1/9

1	MSVDEICENTKMGREYALLGNYETSLVYYQGVLQQIQKLL
41	TSVHEPQRKHQWQTIRQELSQEYEHVKNITKTLNGFKSEP
81	<u>AAPEPAPNGRAAPFSHHOHA</u> AKPAAAEPARDPDVWPPPTP
121	VDHRPSPPYQRAARK <u>DPPRRSEPSKPANRAPG</u> NDRGGRGP
161	${\tt SDRRGDARSGGGRGGARGSDKDKNRGGKSDKDKKAPSGE}$
201	EGDEKKFDPAGYDKDLVENLERDIVORNPNVHWADIAGLT
241	<u>EAKRLLEEAVVLPLWMPDYFK</u> GIRRPWKGVLMV <mark>GPPGTGK</mark>
281	${\tt \ref{thm:lambda}} {\tt \r$
321	MARFYAPSTIFIDEIDSICSKRGTGSEHEASRRV <u>KSELLI</u>
361	<u>OMDGVSGPSAGEE</u> SSKMVMVLAATNFPWDIDEALRRRLEK
401	RIYIPLPEIDGREQLLRINLKEVPLADDIDLKSIAEKMDG
441	YSGA <u>DITNVCRDASMMAM</u> RRR <u>IOGLR</u> PEEIRHIPKEELNQ
481	PSTPADFLLALQKVSKSVGKEDLVKYMAWMEEFGSV

FIG. 1A.

VHWADIAGLTEAKRLLEBAVVLPLWMPDYFKGIFFPW.KGVLMVGPPGTG MSLDDIIGMHDVKGVLHEAVTLPLLVPEFFQGLRSPW.KGMVLAGPPGTG STYDMVGGLTKQIKEIKEVIELBVKHPEIFESLGIAQPKGVILYGPPGTG TTFADVAGCDEAKEEVAELVEY.LREBSRFQKLGGKEPKGVLMVGPPGTG IKWGDIGALANAKDVLLETLEWÐTKYEPÍEVNCPLRLRSGILLIYGYPGGG EKMGIGGLDKEFSDIFRRAFASRVFPÐEIVEQMGCKHVKGILLIYGPPGCG	KTMLAKAVATECGTTEFN.VSSASLTSKYHGESEKLVRLLEEMARFYAP. KTITARAVAHHTDCKFIR.VSSTDLSSKWRGDSEKIVRLLEELARFYAP. KTLLARAVAHHTDCKFIR.VSGAELVOKYIGEGSRAVRELFVAAREHAP. KTLLAKAIAGEAKVPFFT.ISGSDFVEMFVGVGASRVRDMFEQAKKAAP. KTLLASAVAQOCGLNEIS.VKGPEILNKFIGASEONIRELFERAQSVKE.	STIFIDEIDSJCSKRGTGSEHEASRRVKSELLIQMDGVSGPSIIFIDEIDTLGGGRGNSGEHEASRRVKSEFLVQMDG SIIFMDEIDSJGSTRVEGSGGDSEVQRTMLELLNGJDGFET CIIFIDEIDAVGRORGAGLGGGHDEREQTLNQMLVEMDGFES CIIFFDEEDSJAPKRGHDSTGVTDRVVNQLITQMDGAEGL	SAGEESSKMVMVLAATNFPWDIDEALRR. RLEKRIYIPLPEIDGREQLL SONKFDSRRVFVLAATNIPWELDEALRR. RFEKRIFIPLPDIDARKKLI SONKFDSRRVFVLAATRIDIIDPALLRPGRIDRKIEFPPPSVAARAETI N. GVYILAATGRPDLIDBALLRPGRFDRQVVVGLPDVRGREQIL N. NILVIGMTNRPDLIDSALLRPGRLDKSVICNIPTESERLDIL	RINLKEVPEJADDĪĒJKS IĀEKMDGĶSGADĪT EKSMEGTPKSDEĪNYDDLAARĪJEGFSGADVV RIHSRKMNĒJTRGĪNLRKVAEKMNGŪSGADVK KVHMRRVPIJAPDĪDĀAI IĀRGĀPGFSFADĒLA QAIVNSKDKDTGQKKFALEKNADĪKLIĀEKĀPĀGFSFADĒLQ HIHTARMRGHQLĒSADVDĪKELĪĀVĒĪKNĒSĒĀĒLE
231 190 145 149 224	280 239 195 198 744	328 287 246 792 324	369 324 285 288 373	417 372 329 332 875
p60 mei-1 Sug1p ftsH PAS1p NSF	p60 mei-1 Sug1p ftsH Pas1p NSF	p60 mei-1 Suglp ftsH Paslp	p60 mei-1 Suglp ftsH Paslp	p60 mei-1 Sug1p ftsH Pas1p NSF

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Fig. 1A
                                        TSYHEPQRKHQWQTIRQELSQEYEHYKNITKTLNGFKSEP
                             4 1
                                         A A P E P A P N H R A A P F S H H Q H A A K P A A A E P A R D P D Y W P P P T P
                             8 1
                                        YD HRP SPPYQRAARK DPPRRSEPSKPANRAPGNDRGGRGP
                             121
                                         SDRRGDARSGGGRGGARGSDKDKNRGGKSDKDKKAPSGE
                             161
                                        EGDEKKFDPAGYDKDLVENLERDIYQRNPNYHWADIAGLT
                             201
                                        EAKRLL EEAVVLPLWMPDYFKGIRRPWKGYLMYGPPGTGK
                             241
                                        THE LAKAVATECGTTFFN VSSASLTSKYHGE SEKL VRLLFE
                             281
                                        MARFYAPSTIFIDEIDSICSKRGTGSEHEASRRYKSELLI
                             321
                                         QMDGYSGPSAGEESSKMYMYLAATNFPWDIDEALRRRLEK
                             361
                                         RIYIPLPEIDGREQLLRINLKEYPLADDIDLKSIAEKMDG
                             401
                                         Y S G A D I T N Y C R D A S M M A M R R R I Q G L R P E E I R H I P K E E L N Q
                             441
                                         P S T P A D F L L A L Q K Y S K S Y G K E D L Y K Y M A W M E E F G S Y
                              481
                                              VHWADIAGLTEAKRLLEBAYYLELWMEDYBKGIRRPW.KGYMYSPPGHG
MSLDDIIGMHDYBQYLHBAYTLELLYEEFBQGLRSPW.KAMYLASEPGHG
STYDMYGGLTKQIKEIKBYIELEYKHBELBESLGIAQPKGYILYSEPGHS
TTFABYAGCDEAKEEYABLYEY.LREESRBQKLGGKIPKGYEMYSEPGHG
IKWGDIGALANARDYLLETLEWETKYEPIEYNCPLRLRSGIELYSYPGCS
EKMGIGGLDKEFSDIFRRAFASRYFPEEIVEQMGCKHYKGIELYSEPGCS
                                    231
 Fig. 1B
                        p60
                                    190
                                    145
                        Sug 1p
                        ftsH
                                     149
                        Pas1p
                                    694
                                     224
                        NSF
                                              KTMBAKAYATECGTTEFN. VSSASLTSKYHSESEKLVRLEREMARFYAE
KTRIARAIASESSSTEFT. VSSTDLSSKVRSDSEKIVRLERELARFYAD
KTABARAYAHHTDCKEIR. WSGAELYQKYISEGSRMVREID VMAREHAD
KTALAKAIAGEAKYPEFT. ISGSDFYEMFYSYGASRVRDMEEQAKKAMD
KTABASAYAQQCGLMEIS. YKGPEILMKFISASEQNIREMEERAQSYKD
KTEDARQIGKMLNAREPKYVNAPEILMKYVSESEANIRKMEADAEEEQRR
                       p60
                                     280
                        mei-1
                                     239
                                    195
                        Sug1p
                                     198
                        ftsH
                        Pas1p
                                     744
                        NSF
                                     274
                                              STIE IDE ED SIC SKR... GTGSE HE ASRRYKSE IN LONDGYSGP
SE IE IDE ED TLGGOR... EN SGE HE ASRRYKSE IN LONDG...
SE IE IDE ED SIGSTRYE GS GGDSE VORTHLE IN NOL DGF ET.
CILE IDE ED AVGRORGA GLGGHDERE OTLNOM VEMDGE GG... DS TG YTD EV YNOLL TOMOGAEGL
LGANSGLHELIF DE ED ATCKORGS. NAGSTGYHD TV YNOLL SKIDG VEOL
                        p60
                                     328
                        mei-1
                                     287
                        Sug1p
                                     243
                        ftsH
                                     246
                        Pas1p
                                     792
                        NSF
                                              SAGEESSKMYMYLNATNEEWDIDE ALRR. RLEKRIYIPLEEIDGREQLE
SQNKFDSRYFWLMATNIEWELDEALRR. RFEKRIFIPEDDIDARKKLI
SKNIKIIMATNREDILDEALRR. RFEKRIFIPEDDIDARKKLI
NEGIIWIAATNREDVLDEALRREGREDRQYYYGEPDYRGREQIE
D. GYYILAATSREDLIDSALIREGREDKSVICNIETESERLDIE
N. NILWIGHTNREDLIDE ALRREGRELDKSVICNIETESERLDIE
                        p60
                                     369
                        mei-1
                                     324
                        Sug 1p
                                     285
                        ftsH
                                     288
                        Pas1p
                                     832
                        NSF
                                              RINL KEYPIADDIDLKSIAEKMDGYSGAPIT
EKSM EGTPK SDEINYDDLAARMEGESGAD YY
RIHS RKMNITRGINLRK VAEKMNGCSGAD VK
KYHM RRYPIAPDIDAAIIARGIPGESGADLA
QAI YNSKD KDTGQ KKF AI EKNADLKLIAEKIJAGE SGADLQ
HI HTARMRGHQ LISAD VDIKELAY EN KNESGAELE
                        p60
                                     417
                        mei-1
                                     372
                                     329
                         Sug1p
                         ftsH
                                     332
                        Pasip
                                     875
                        NSF
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1

MSYDEICENTKMGREYALLGNYETSLYYYQGYLQQIQKLL